## SEQUENCE LISTING

<110> Universitätsklinikum Freiburg

<120> THE PRV-1 GENE AND USE THEREOF

<130> E980930

<140> PCT/EP00/09594

<141> 2000-09-29

<150> DE 199 47 010.3

<151> 1999-09-30

<160> 10

<170> PADAT Sequenzmodul, Version 1.0

<210> 1

<211> 1600

<212> DNA

<213> homo sapiens

<220>

<223>

<400> 1

aaaagcagaa agagattacc	agccacagac	gggtcatgag	cgcggtatta	ctgctggccc	60
tcctggggtt catcctccca	ctgccaggag	tgcaggcgct	gctctgccag	tttgggacag	120
ttcagcatgt gtggaaggtg	tccgacctgc	cccggcaatg	gacccctaag	aacaccagct	180
gcgacagcgg cttggggtgc	caggacacgt	tgatgctcat	tgagagcgga	ccccaagtga	240
gcctggtgct ctccaagggc	tgcacggagg	ccaaggacca	ggagccccgc	gtcactgagc	300
accggatggg ccccggcctc	tocotgatot	cctacacctt	cgtgtgccgc	caggaggact	360
tctgcaacaa cctcgttaac	tccctcccgc	tttgggcccc	acagccccca	gcagacccag	420
gatccttgag gtgcccagtc	tgcttgtcta	tggaaggctg	tctggagggg	acaacagaag	480
agatctgccc caaggggacc	acacactgtt	atgatggcct	cctcaggctc	aggggaggag	540
gcatcttctc caatctgaga	gtccagggat	gcatgcccca	gccaggttgc	aacctgctca	600
atgggacaca ggaaattggg	cccgtgggta	tgactgagaa	ctgcaatagg	aaagattttc	660
tgacctgtca tcgggggacc	accattatga	cacacggaaa	cttggctcaa	gaacccactg	720
attggaccac atcgaatacc	gagatgtgcg	aggtggggca	ggtgtgtcag	gagacgctgc	780
tgctcataga tgtaggactc	acatcaaccc	tggtggggac	aaaaggctgc	agcactgttg	840

ttcccagaag	accaccatcc	actcagcccc	tcctggggtg	cttgtggcct	900
cttctgctcc	tcggacctgt	gcaatagtgc	cagcagcagc	agcgttctgc	960
ccctcctcaa	gctgcccctg	tcccaggaga	ccggcagtgt	cctacctgtg	1020
tggaacctgt	tcaagtggct	cccccgaat	gacctgcccc	aggggcgcca	1080
tgatgggtac	attcatctct	caggaggtgg	gctgtccacc	aaaatgagca	1140
					1200
					1260
					1320
					1380
					1440
					1500
					1560
					1600
	cttctgctcc ccctcctcaa tggaacctgt tgatgggtac cgtggcccaa gcgtgagaag cctggagtct ggtttgccct cactcaacct ctgtccatga ctggggagag	cttctgctcc tcggacctgt ccctcctaa gctgcccctg tggaacctgt tcaagtggct tgatgggtac attcatctct cgtggcccaa ccttccagct gcgtgagaag cgtgatgtgc cctggagtct ctcacttggg ggtttgccct tcctgctaac cactcaacct ccctctgacc ctgtccatga atcatcttcc ctggggagag cctggagcat	cttctgctcc tcggacctgt gcaatagtgc ccctccteaa gctgcccctg tcccaggaga tggaacctgt tcaagtggct cccccgaat tgatgggtac attcatctct caggaggtgg cgtggccaa ccttccagct tcttgttgaa gcgtgagaag cgtgatgtgc agcctcctgc cctggagtct ctcacttggg gggtggggct ggtttgccct tcctgctaac tctattaccc cactcaacct ccctctgacc tcataaccta ctgtccatga atcatcttcc ccacacaca	cettetgetee teggacetgt geaatagtge cageageage cecteeteaa getgeecetg teceaggaga eeggeagtgt tggaacetgt teaagtgget eegegagtgg getgteeaee egtggeecaa eetteeaget tettgttgaa eeaeaeaga gegtgagaaag egtgatgtge ageeteetge eteteageat eeteggagtet eteetgegggget gggtgggget ggeaetggee ggtttgeeet teetgetaae tettataeee eeaegatet eetetgaee teetgaee teetgeee eetetgee eetetgeggagag eeteggageat eeggaeettge eetatgggag	tteccagaag accaceatec acteagecec tectggggtg cttggcect tectgectec teggacetgt geaatagtge cageageage agegttetge eccetecteaa getggecetg teccaggaga eeggeagtgt cetacetgtg tggaacetgt teaagtgget cageaggtgg getgtecace aggggegeagtg eggtgggeceaa ecttecaget tettgtgaa ecaceagaa caaateggga geggtggagaag eggtgatgtg ageetteetge etctetge ggcaetgge eccagaateggggggggggggggggggggggggggggggg

<210> 2

<211> 437

<212> PRT

<213> homo sapiens

<400> 2

Met Ser Ala Val Leu Leu Ala Leu Leu Gly Phe Ile Leu Pro Leu 10 Pro Gly Val Gln Ala Leu Leu Cys Gln Phe Gly Thr Val Gln His Val 25 Trp Lys Val Ser Asp Leu Pro Arg Gln Trp Thr Pro Lys Asn Thr Ser Cys Asp Ser Gly Leu Gly Cys Gln Asp Thr Leu Met Leu Ile Glu Ser 60 55 Gly Pro Gln Val Ser Leu Val Leu Ser Lys Gly Cys Thr Glu Ala Lys 75 70 Asp Gln Glu Pro Arg Val Thr Glu His Arg Met Gly Pro Gly Leu Ser 85 90 Leu Ile Ser Tyr Thr Phe Val Cys Arg Gln Glu Asp Phe Cys Asn Asn 105 110 100 Leu Val Asn Ser Leu Pro Leu Trp Ala Pro Gln Pro Pro Ala Asp Pro 125 120 Gly Ser Leu Arg Cys Pro Val Cys Leu Ser Met Glu Gly Cys Leu Glu 135 130 Gly Thr Thr Glu Glu Ile Cys Pro Lys Gly Thr Thr His Cys Tyr Asp 155 150 Gly Leu Leu Arg Leu Arg Gly Gly Gly Ile Phe Ser Asn Leu Arg Val 175 170 165

3

```
Gln Gly Cys Met Pro Gln Pro Gly Cys Asn Leu Leu Asn Gly Thr Gln
                                185
Glu Ile Gly Pro Val Gly Met Thr Glu Asn Cys Asn Arg Lys Asp Phe
                                                 205
                            200
        195
Leu Thr Cys His Arg Gly Thr Thr Ile Met Thr His Gly Asn Leu Ala
                        215
Gln Glu Pro Thr Asp Trp Thr Thr Ser Asn Thr Glu Met Cys Glu Val
                                        235
                    230
Gly Gln Val Cys Gln Glu Thr Leu Leu Leu Ile Asp Val Gly Leu Thr
                                    250
                245
Ser Thr Leu Val Gly Thr Lys Gly Cys Ser Thr Val Gly Ala Gln Asn
                                265
            260
Ser Gln Lys Thr Thr Ile His Ser Ala Pro Pro Gly Val Leu Val Ala
                                                 285
                            280
Ser Tyr Thr His Phe Cys Ser Ser Asp Leu Cys Asn Ser Ala Ser Ser
                                             300
Ser Ser Val Leu Leu Asn Ser Leu Pro Pro Gln Ala Ala Pro Val Pro
                                         315
                    310
Gly Asp Arg Gln Cys Pro Thr Cys Val Gln Pro Leu Gly Thr Cys Ser
                                     330
Ser Gly Ser Pro Arg Met Thr Cys Pro Arg Gly Ala Thr His Cys Tyr
                                345
            340
Asp Gly Tyr Ile His Leu Ser Gly Gly Gly Leu Ser Thr Lys Met Ser
                                                 365
                            360
        355
Ile Gln Gly Cys Val Ala Gln Pro Ser Ser Phe Leu Leu Asn His Thr
                                             380
                        375
Arg Gln Ile Gly Ile Phe Ser Ala Arg Glu Lys Arg Asp Val Gln Pro
                                         395
                    390
Pro Ala Ser Gln His Glu Gly Gly Gly Ala Glu Gly Leu Glu Ser Leu
                405
                                    410
Thr Trp Gly Val Gly Leu Ala Leu Ala Pro Ala Leu Trp Trp Gly Val
                                                     430
                                 425
            420
Val Cys Pro Ser Cys
        435
<210> 3
<211> 24
<212> RNA
<213> Artificial Sequence
<220>
<223> 5'-end of PRV-1-sequence
<400> 3
```

24

<210> 4 <211> 24 <212> RNA <213> Artificial Sequence <220> <223> Antisense-Molecule

aaaagcagaa agagattacc agcc

4

```
<400> 4
                                                               24
ggctggtaat ctctttctgc tttt
<210> 5
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> amino acids 34-46 of PRV-1
<400> 5
Lys Val Ser Asp Leu Pro Arg Gln Trp Thr Pro Lys Asn
<210> 6
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> amino acids 391-405 of PRV-1
<400> 6
Ser Ala Arg Glu Lys Arg Asp Val Gln Pro Pro Ala Ser Gln His
  1
<210> 7
<211> 27
<212> DNA
<213> Artificial Sequence
<220>
<223> RT-Primer
<400> 7
                                                               27
attaggttat gaggtcagag ggaggtt
<210> 8
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
```

<223> sense-Primer

<400> 8	
gcagaaagag attaccagcc acagacgg	28
<210> 9 <211> 28 <212> DNA <213> Artificial Sequence	
<220> <223> antisense-Primer	
<400> 9	
gaatcgtggg ggtaatagag ttagcagg	28
<210> 10 <211> 29 <212> DNA <213> Artificial Sequence	
<220> <223> probe	
<400> 10	
ttettgttga accaecag acaaategg	29